



IFWO

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/10/825,898

DATE: 09/01/2004  
 TIME: 13:06:27

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1 <110> APPLICANT: BOYLE, WILLIAM  
 2 <120> TITLE OF INVENTION: OSTEOPROTEGERIN BINDING PROTEINS AND RECEPTORS  
 3 <130> FILE REFERENCE: A-451K REV 09-10-03 54SEQ  
 C--> 4 <140> CURRENT APPLICATION NUMBER: US/10/825,898  
 C--> 5 <141> CURRENT FILING DATE: 2004-04-15  
 6 <150> PRIOR APPLICATION NUMBER: US/10/825,898  
 7 <151> PRIOR FILING DATE: 2004-04-15  
 8 <150> PRIOR APPLICATION NUMBER: US 09/052,521  
 9 <151> PRIOR FILING DATE: 1998-03-30  
 10 <150> PRIOR APPLICATION NUMBER: US 08/880,855  
 11 <151> PRIOR FILING DATE: 1997-06-23  
 12 <150> PRIOR APPLICATION NUMBER: US 08/842,842  
 13 <151> PRIOR FILING DATE: 1997-04-16  
 14 <160> NUMBER OF SEQ ID NOS: 54  
 15 <170> SOFTWARE: PatentIn version 3.1  
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 18 <211> LENGTH: 2295  
 19 <212> TYPE: DNA  
 20 <213> ORGANISM: Mus musculus  
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 23 <222> LOCATION: (158)..(1105)  
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 28 tcgcggagca gggcgcccga actccgggag ccgcgcc atg cgc cgg gcc agc cga 175  
 29 Met Arg Arg Ala Ser Arg  
 30 1 5  
 31 gac tac ggc aag tac ctg cgc agc tcg gag gag atg ggc agc ggc ccc 223  
 32 Asp Tyr Gly Lys Tyr Leu Arg Ser Ser Glu Glu Met Gly Ser Gly Pro  
 33 10 15 20  
 34 ggc gtc cca cac gag ggt ccg ctg cac ccc gcg cct tct gca ccg gct 271  
 35 Gly Val Pro His Glu Gly Pro Leu His Pro Ala Pro Ser Ala Pro Ala  
 36 25 30 35  
 37 ccg gcg ccg cca ccc gcc gcc tcc cgc tcc atg ttc ctg gcc ctc ctg 319  
 38 Pro Ala Pro Pro Pro Ala Ala Ser Arg Ser Met Phe Leu Ala Leu Leu  
 39 40 45 50  
 40 ggg ctg gga ctg ggc cag gtg gtc tgc agc atc gct ctg ttc ctg tac 367  
 41 Gly Leu Gly Leu Gly Gln Val Val Cys Ser Ile Ala Leu Phe Leu Tyr  
 42 55 60 65 70  
 43 ttt cga gcg cag atg gat cct aac aga ata tca gaa gac agc act cac 415  
 44 Phe Arg Ala Gln Met Asp Pro Asn Arg Ile Ser Glu Asp Ser Thr His

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47	Cys Phe Tyr Arg Ile Leu Arg Leu His Glu Asn Ala Gly Leu Gln Asp						
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49	tcg act ctg gag agt gaa gac aca cta cct gac tcc tgc agg agg atg						511
50	Ser Thr Leu Glu Ser Glu Asp Thr Leu Pro Asp Ser Cys Arg Arg Met						
51		105		110		115	
52	aaa caa gcc ttt cag ggg gcc gtg cag aag gaa ctg caa cac att gtg						559
53	Lys Gln Ala Phe Gln Gly Ala Val Gln Lys Glu Leu Gln His Ile Val						
54		120		125		130	
55	ggg cca cag cgc ttc tca gga gct cca gct atg atg gaa ggc tca tgg						607
56	Gly Pro Gln Arg Phe Ser Gly Ala Pro Ala Met Met Glu Gly Ser Trp						
57		135		140		145	
58	ttg gat gtg gcc cag cga ggc aag cct gag gcc cag cca ttt gca cac						655
59	Leu Asp Val Ala Gln Arg Gly Lys Pro Glu Ala Gln Pro Phe Ala His						
60		155		160		165	
61	ctc acc atc aat gct gcc agc atc cca tcg ggt tcc cat aaa gtc act						703
62	Leu Thr Ile Asn Ala Ala Ser Ile Pro Ser Gly Ser His Lys Val Thr						
63		170		175		180	
64	ctg tcc tct tgg tac cac gat cga ggc tgg gcc aag atc tct aac atg						751
65	Leu Ser Ser Trp Tyr His Asp Arg Gly Trp Ala Lys Ile Ser Asn Met						
66		185		190		195	
67	acg tta agc aac gga aaa cta agg gtt aac caa gat ggc ttc tat tac						799
68	Thr Leu Ser Asn Gly Lys Leu Arg Val Asn Gln Asp Gly Phe Tyr Tyr						
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70	ctg tac gcc aac att tgc ttt cgg cat cat gaa aca tcg gga agc gta						847
71	Leu Tyr Ala Asn Ile Cys Phe Arg His His Glu Thr Ser Gly Ser Val						
72		215		220		225	
73	cct aca gac tat ctt cag ctg atg gtg tat gtc gtt aaa acc agc atc						895
74	Pro Thr Asp Tyr Leu Gln Leu Met Val Tyr Val Val Lys Thr Ser Ile						
75		235		240		245	
76	aaa atc cca agt tct cat aac ctg atg aaa gga ggc agc acg aaa aac						943
77	Lys Ile Pro Ser Ser His Asn Leu Met Lys Gly Gly Ser Thr Lys Asn						
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79	tgg tcg ggc aat tct gaa ttc cac ttt tat tcc ata aat gtt ggc gga						991
80	Trp Ser Gly Asn Ser Glu Phe His Phe Tyr Ser Ile Asn Val Gly Gly						
81		265		270		275	
82	ttt ttc aag ctc cga gct ggt gaa gaa att agc att cag gtg tcc aac						1039
83	Phe Phe Lys Leu Arg Ala Gly Glu Glu Ile Ser Ile Gln Val Ser Asn						
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85	cct tcc ctg ctg gat ccg gat caa gat gcg acg tac ttt ggc gct ttc						1087
86	Pro Ser Leu Leu Asp Pro Asp Gln Asp Ala Thr Tyr Phe Gly Ala Phe						
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89	Lys Val Gln Asp Ile Asp						
90		315					
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92	actaagagac atggcccacg gtgtatgaaa ctccacagccc tctctcttga gcctgtacag						1255
93	gttgtgtata tgtaaagtcc ataggtgatg ttagattcat ggtgattaca caacggtttt						1315

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97  tgaaggggta agttcttttg aattgtttaca ttgcgctggg acctgcaaat aagttctttt 1555
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101 actggtgcac tttgtaattc ccctgaaggt actcgtagct aagggggcag aatactgttt 1795
102 ctggtgacca catgtagttt atttctttat tctttttaac ttaatagagt cttcagactt 1855
103 gtcaaaacta tgcaagcaaa ataaataaat aaaaataaaa tgaatacctt gaataataag 1915
104 taggatgttg gtcaccaggt gcctttcaaa tttagaagct aattgacttt aggagctgac 1975
105 atagccaaaa aggatacata ataggctact gaaatctgtc aggagtattt atgcaattat 2035
106 tgaacagggtg tcttttttta caagagctac aaattgtaaa ttttgtttct ttttttccc 2095
107 atagaaaatg tactatagtt tatcagccaa aaaacaatcc actttttaat ttagtgaaag 2155
108 ttattttatt atactgtaca ataaaagcat tgtctctgaa tggttaatttt ttggtacaaa 2215
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122 35 40 45
123 Met Phe Leu Ala Leu Leu Gly Leu Gly Leu Gly Gln Val Val Cys Ser
124 50 55 60
125 Ile Ala Leu Phe Leu Tyr Phe Arg Ala Gln Met Asp Pro Asn Arg Ile
126 65 70 75 80
127 Ser Glu Asp Ser Thr His Cys Phe Tyr Arg Ile Leu Arg Leu His Glu
128 85 90 95
129 Asn Ala Gly Leu Gln Asp Ser Thr Leu Glu Ser Glu Asp Thr Leu Pro
130 100 105 110
131 Asp Ser Cys Arg Arg Met Lys Gln Ala Phe Gln Gly Ala Val Gln Lys
132 115 120 125
133 Glu Leu Gln His Ile Val Gly Pro Gln Arg Phe Ser Gly Ala Pro Ala
134 130 135 140
135 Met Met Glu Gly Ser Trp Leu Asp Val Ala Gln Arg Gly Lys Pro Glu
136 145 150 155 160
137 Ala Gln Pro Phe Ala His Leu Thr Ile Asn Ala Ala Ser Ile Pro Ser
138 165 170 175
139 Gly Ser His Lys Val Thr Leu Ser Ser Trp Tyr His Asp Arg Gly Trp
140 180 185 190
141 Ala Lys Ile Ser Asn Met Thr Leu Ser Asn Gly Lys Leu Arg Val Asn
142 195 200 205
143 Gln Asp Gly Phe Tyr Tyr Leu Tyr Ala Asn Ile Cys Phe Arg His His

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149      Gly Gly Ser Thr Lys Asn Trp Ser Gly Asn Ser Glu Phe His Phe Tyr
150          260          265          270
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152          275          280          285
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170      cgcc atg cgc cgc gcc agc aga gac tac acc aag tac ctg cgt ggc tcg      229
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174      Glu Glu Met Gly Gly Gly Pro Gly Ala Pro His Glu Gly Pro Leu His
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176      gcc ccg ccg ccg cct gcg ccg cac cag ccc ccc gcc gcc tcc cgc tcc      325
177      Ala Pro Pro Pro Pro Ala Pro His Gln Pro Pro Ala Ala Ser Arg Ser
178          35          40          45
179      atg ttc gtg gcc ctc ctg ggg ctg ggg ctg ggc cag gtt gtc tgc agc      373
180      Met Phe Val Ala Leu Leu Gly Leu Gly Leu Gly Gln Val Val Cys Ser
181          50          55          60
182      gtc gcc ctg ttc ttc tat ttc aga gcg cag atg gat cct aat aga ata      421
183      Val Ala Leu Phe Phe Tyr Phe Arg Ala Gln Met Asp Pro Asn Arg Ile
184          65          70          75
185      tca gaa gat ggc act cac tgc att tat aga att ttg aga ctc cat gaa      469
186      Ser Glu Asp Gly Thr His Cys Ile Tyr Arg Ile Leu Arg Leu His Glu
187      80          85          90          95
188      aat gca gat ttt caa gac aca act ctg gag agt caa gat aca aaa tta      517
189      Asn Ala Asp Phe Gln Asp Thr Thr Leu Glu Ser Gln Asp Thr Lys Leu
190          100          105          110
191      ata cct gat tca tgt agg aga att aaa cag gcc ttt caa gga gct gtg      565
192      Ile Pro Asp Ser Cys Arg Arg Ile Lys Gln Ala Phe Gln Gly Ala Val
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197	aaa gcg atg gtg gat ggc tca tgg tta gat ctg gcc aag agg agc aag	661
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238	agttatatatt cagatgtaat gttttctttg caaagtattg taaattatat ttgtgctata	1675
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242	aatagatttt ttcagacttg tcaagcctgt gcaaaaaaat taaaatggat gccttgaata	1915

VERIFICATION SUMMARY

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Output Set: N:\CRF4\09012004\J825898.raw

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:5 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
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